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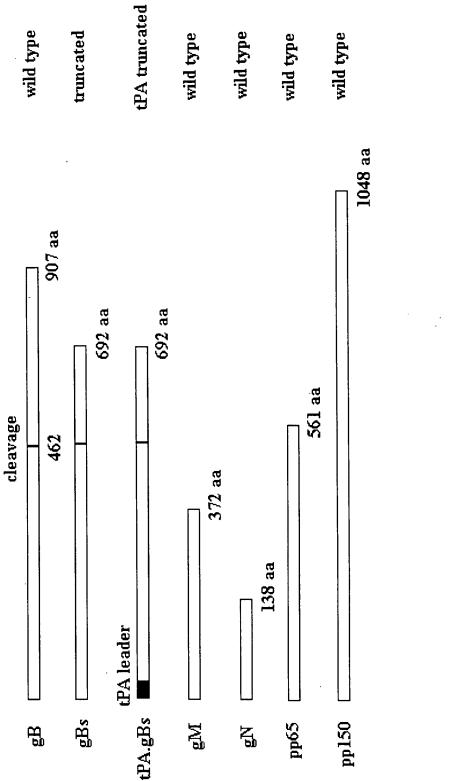
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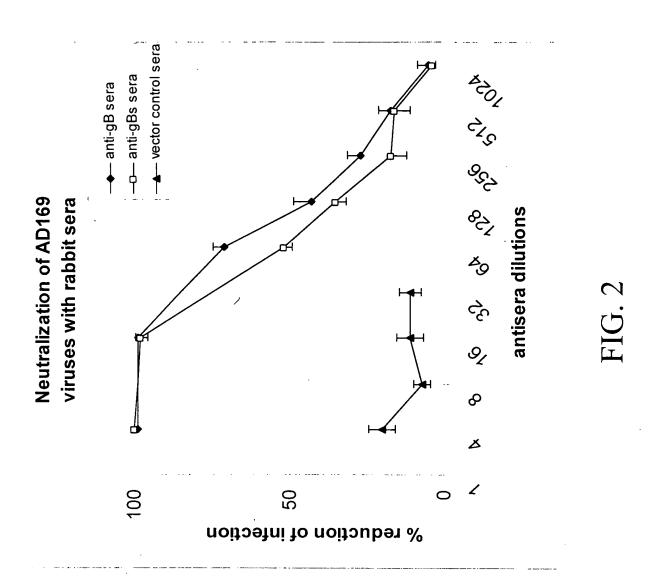
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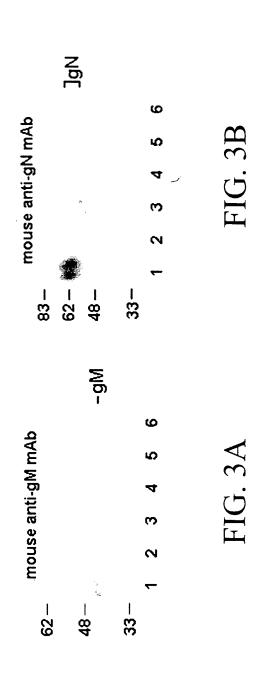
Matter No.: 07917-190001 Page 1 of 20 Applicant(s): Lu et al.
COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS TREATMENT



Matter No.: 07917-190001 Page 2 of 20 Applicant(s): Lu et al. COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS TREATMENT

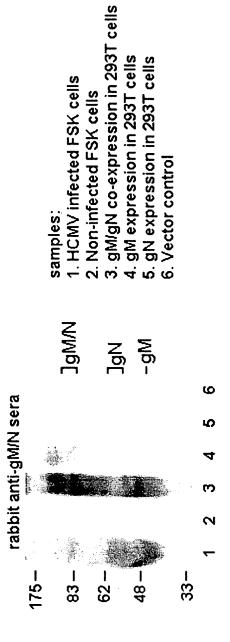


Applicant(s): Lu et al.
COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS
TREATMENT



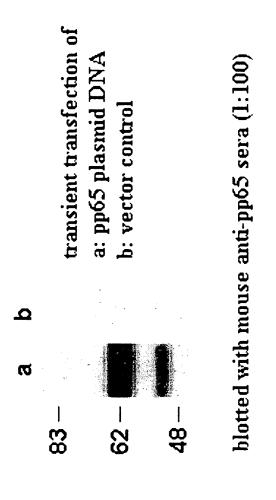
Matter No.: 07917-190001 Page 4 of 20 Applicant(s): Lu et al. COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

TREATMENT

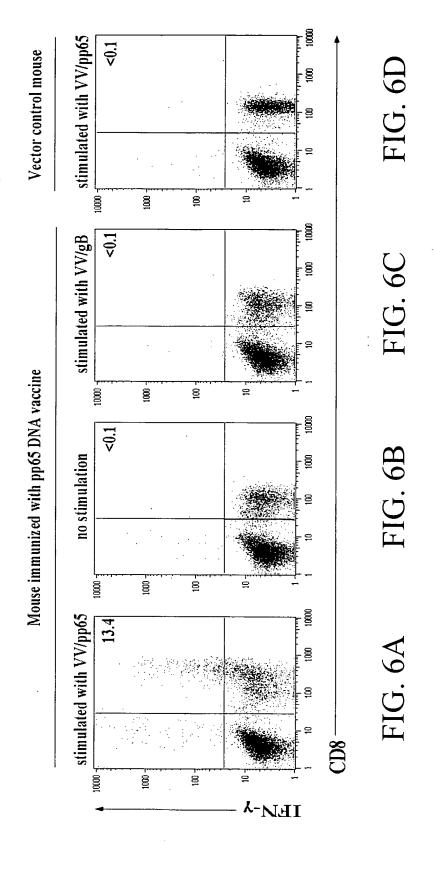


Matter No.: 07917-190001 Page 5 of 20 Applicant(s): Lu et al. COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

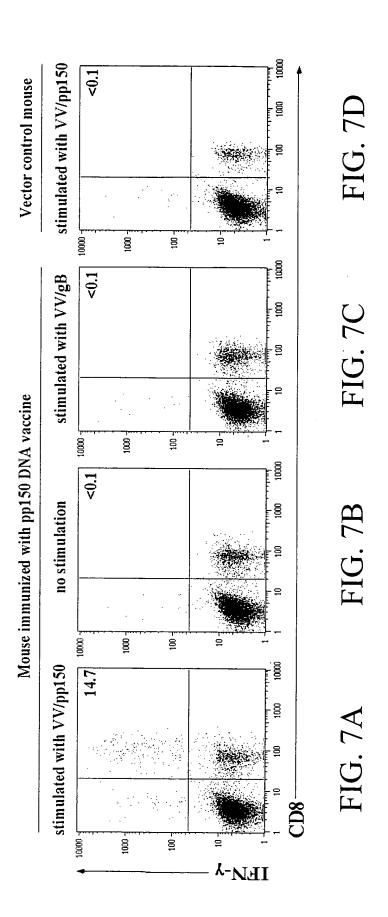
TREATMENT

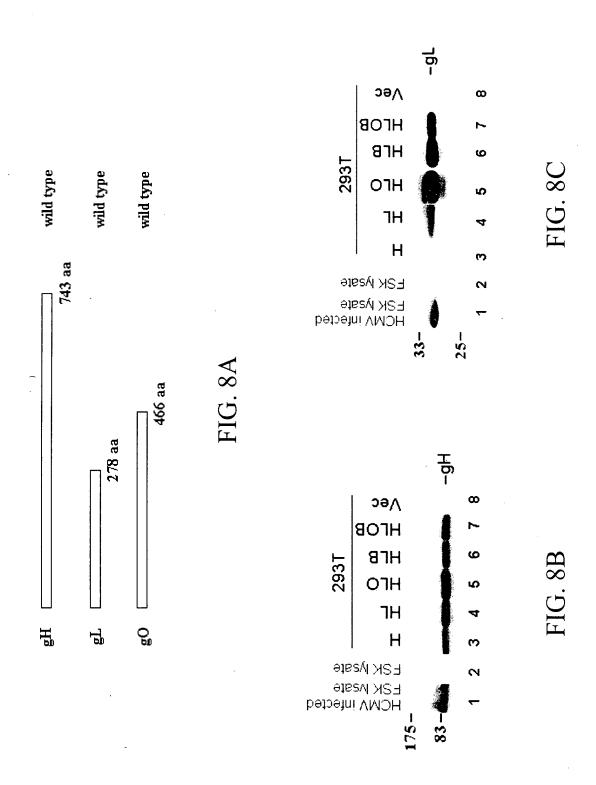


Matter No.: 07917-190001 Page 6 of 20 Applicant(s): Lu et al. COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS TREATMENT

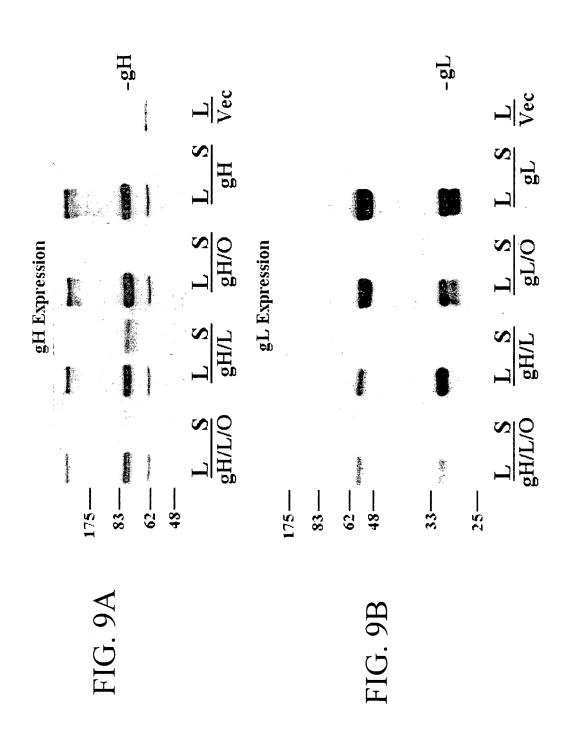


Matter No.: 07917-190001 Page 7 of 20 Applicant(s): Lu et al. COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS TREATMENT





Matter No.: 07917-190001 Page 9 of 20 Applicant(s): Lu et al. COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS TREATMENT



Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

Page 10 of 20

TREATMENT

HCMV gB cDNA Sequence

CGGTTTCCTCTTCTAGTACTTCCCATGCAACTTCTTCTACTCACAATGGAAGCCATACTTCTCG TACGACGTCTGCTCAAACCCGGTCAGTCTATTCTCAACACGTAACGTCTTCTGAAGCCGTCAG TCATAGAGCCAACGAGACTATCTACAACACTACCCTCAAGTACGGAGATGTGGTGGGAGTCA ACACTACCAAGTACCCCTATCGCGTGTTCTATGGCCCAGGGTACGGATCTTATTCGCTTTGA ACGTAATATCATCTGCACCTCGATGAAGCCTATCAATGAAGACTTGGATGAGGGCATCATGGT GGTCTACAAGCGCAACATCGTGGCGCACACCTTTAAGGTACGGGTCTACCAAAAGGTTTTGAC GTTTCGTCGTAGCTACGCTTACATCTACACCACTTATCTGCTGGGCAGCAATACGGAATACGT GGCGCCTCCTATGTGGGAGATTCATCACATCAACAAGTTTGCTCAATGCTACAGTTCCTACAG CCGCGTTATAGGAGGCACGGTTTTCGTGGCATATCATAGGGACAGTTATGAAAACAAAACCAT GCAATTAATTCCCGACGATTATTCCAACACCCACAGTACCCGTTACGTGACGGTCAAGGATCA GTGGCACAGCCGCGGCAGCACCTGGCTCTATCGTGAGACCTGTAATCTGAACTGTATGCTGAC CATCACTACTGCGCGCTCCAAGTATCCTTATCATTTTTTTGCAACTTCCACGGGTGATGTGGTT TACATTTCTCCTTTCTACAACGGAACCAATCGCAATGCCAGCTACTTTGGAGAAAACGCCGAC AAGTTTTCATTTTCCCGAACTACACCATCGTTTCCGACTTTGGAAGACCCAACGCTGCGCCAG AAACCCATAGGTTGGTGGCTTTTCTCGAACGTGCCGACTCGGTGATCTCTTGGGATATACAGG ACGAGAÁGAATGTCACCTGCCAGCTCACCTTCTGGGAAGCCTCGGAACGTACTATCCGTTCCG AAGCCGAAGACTCGTACCACTTTTCTTCTGCCAAAATGACTGCAACTTTTCTGTCTAAGAAAC CAGCAGATTTTCAATACTTCATACAATCAAACATATGAAAAATACGGAAACGTGTCCGTCTTC GAAACCAGCGGCGTCTGGTGGTGTTCTGGCAAGGCATCAAGCAAAAATCTTTGGTGGAATT GGAACGTTTGGCCAATCGATCCAGTCTGAATATCACTCATAGGACCAGAAGAAGTACGAGTG ACAATAATACAACTCATTTGTCCAGCATGGAATCGGTGCACAATCTGGTCTACGCCCAGCTGC AGTTCACCTATGACACGTTGCGCGGTTACATCAACCGGGCGCTGGCGCAAATCGCAGAAGCCT GGTGTGTGGATCAACGGCGCACCCTAGAGGTCTTCAAGGAACTCAGCAAGATCAACCCGTCA GCCATTCTCTCGGCCATTTACAACAAACCGATTGCCGCGCGTTTCATGGGTGATGTCTTGGGCC TGGCCAGCTGCGTGACCATCAACCAACCAGCGTCAAGGTGCTGCGTGATATGAACGTGAAG GAATCGCCAGGACGCTGCTACTCACGACCCGTGGTCATCTTTAATTTCGCCAACAGCTCGTAC GTGCAGTACGGTCAACTGGGCGAGGACAACGAAATCCTGTTGGGCAACCACCGCACTGAGGA ATGTCAGCTTCCCAGCCTCAAGATCTTCATCGCCGGGAACTCGGCCTACGAGTACGTGGACTA CCTCTTCAAACGCATGATTGACCTCAGCAGTATCTCCACCGTCGACAGCATGATCGCCCTGGA TATCGACCCGCTGGAAAATACCGACTTCAGGGTACTGGAACTTTACTCGCAGAAAGAGCTGCG TTCCAGCAACGTTTTTGACCTCGAAGAGATCATGCGCGAATTCAACTCGTACAAGCAGCGGGT AAAGTACGTGGAGGACAAGGTAGTCGACCCGCTACCCCAAGGGTCTGGACGACC GCGGTGGCCTCCGTGGTCGAAGGCGTTGCCACCTTCCTCAAAAACCCCTTCGGAGCCTTCACC ATCATCCTCGTGGCCATAGCCGTAGTCATTATCACTTATTTGATCTATACTCGACAGCGGCGTC TGTGCACGCAGCCGCTGCAGAACCTCTTTCCCTATCTGGTGTCCGCCGACGGGACCACCGTGA CGTCGGGCAGCACCAAAGACACGTCGTTACAGGCTCCGCCTTCCTACGAGGAAAGTGTTTATA ATTCTGGTCGCAAAGGACCGGGACCACCGTCGTCTGATGCATCCACGGCGGCTCCGCCTTACA CAGCAGAACGGTACAGATTCTTTGGACGGACAGACTGGCACGCAGGACAAGGGACAGAAGCC TAACCTGCTAGACCGGCTGCGACATCGCAAAAACGGCTACAGACACTTGAAAGACTCCGACG AAGAAGAGAACGTCTGA (SEQ ID NO:1)

Page 11 of 20

Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

TREATMENT

HCMV gB Amino Acid Sequence

MESRIWCLVVCVNLCIVCLGAAVSSSSTSHATSSTHNGSHTSRTTSAQTRSVYSQ HVTSSEAVSHRANETIYNTTLKYGDVVGVNTTKYPYRVCSMAQGTDLIRFERNII CTSMKPINEDLDEGIMVVYKRNIVAHTFKVRVYOKVLTFRRSYAYIYTTYLLGSN TEYVAPPMWEIHHINKFAOCYSSYSRVIGGTVFVAYHRDSYENKTMOLIPDDYS NTHSTRYVTVKDQWHSRGSTWLYRETCNLNCMLTITTARSKYPYHFFATSTGDV VYISPFYNGTNRNASYFGENADKFFIFPNYTIVSDFGRPNAAPETHRLVAFLERAD SVISWDIQDEKNVTCQLTFWEASERTIRSEAEDSYHFSSAKMTATFLSKKQEVNM SDSALDCVRDEAINKLOOIFNTSYNOTYEKYGNVSVFETSGGLVVFWOGIKOKS LVELERLANRSSLNITHRTRRSTSDNNTTHLSSMESVHNLVYAQLQFTYDTLRGY INRALAQIAEAWCVDQRRTLEVFKELSKINPSAILSAIYNKPIAARFMGDVLGLAS CVTINQTSVKVLRDMNVKESPGRCYSRPVVIFNFANSSYVQYGQLGEDNEILLGN HRTEECQLPSLKIFIAGNSAYEYVDYLFKRMIDLSSISTVDSMIALDIDPLENTDFR VLELYSOKELRSSNVFDLEEIMREFNSYKORVKYVEDKVVDPLPPYLKGLDDLM SGLGAAGKAVGVAIGAVGGAVASVVEGVATFLKNPFGAFTIILVAIAVVIITYLIY TRQRRLCTQPLQNLFPYLVSADGTTVTSGSTKDTSLQAPPSYEESVYNSGRKGPG PPSSDASTAAPPYTNEQAYQMLLALARLDAEQRAQQNGTDSLDGQTGTQDKGQ KPNLLDRLRHRKNGYRHLKDSDEEENV* (SEQ ID NO:2)

Matter No.: 07917-190001 Page 12 of 20 Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

TREATMENT

HCMV gM cDNA Sequence

ATGGCCCCTCGCACGTGGATAAGGTGAATACACGGACATGGAGCGCTTCTA TCGTTTTCATGGTGCTGACTTTTGTCAACGTCAGCGTGCATCTAGTGCTGAGC AATTTTCCGCACCTGGGCTACCCCTGCGTCTACTATCACGTCGTGGACTTTGA AAGGCTCAACATGTCGGCCTACAACGTAATGCACCTGCACACGCCTATGCTTT TCTTAGACTCGGTGCAGTTGGTGTGCTACGCCGTGTTCATGCAGCTCGTCTTT TTAGCCGTGACCATCTACTACCTGGTATGCTGGATCAAGATCAGCATGCGCA AGGACAAAGGCATGAGCCTAAACCAGTCGACACGCGACATTTCGTACATGGG CGACAGCCTCACAGCCTTCCTCTTCATTCTCAGCATGGACACGTTCCAACTAT TCACACTGACCATGTCATTTCGGCTGCCCAGCATGATCGCCTTCATGGCCGCC GTGCACTTTTCTGCCTGACCATTTTCAACGTGAGCATGGTCACGCAGTACCG CAGCTACAAACGCTCACTCTTTTCTCCGCGTCTGCACCCCAAGCTCAAAG GTACGGTGCAGTTCCGCACGCTCATCGTCAACCTGGTAGAGGTAGCGCTTGG TTTCAACACCACCGTGGTAGCCATGGCCCTGTGCTACGGCTTCGGAAACAACT TTTTCGTGCGTACAGGCCACATGGTGTTAGCCGTCTTCGTGGTCTACGCTATC ATCTCCATCATCTACTTTTTACTGATCGAGGCCGTCTTTTTTCAATACGTCAAG GTGCAATTCGGCTACCACCTGGGCGCCTTCTTTGGACTCTGCGGCCTCATCTA CCCCATCGTGCAGTACGATACCTTCCTCAGCAACGAATACCGCACCGGCATC AGCTGGTCGTTCGGCATGCTCTTTTTCATATGGGCCATGTTTACGACGTGTCG CGCCGTCCGCTACTTTCGCGGACGCGGTAGCGGCAGTGTCAAGTACCAGGCG CTGGCCACAGCCTCCGGCGAAGAAGTCGCTGTGCTCAGTCACCACGACAGCT TGGAAAGCCGTCGCCTCCGCGAAGAAGAGGACGACGACGATGATGAAGACT TCGAGGACGCTTAA (SEQ ID NO:3)

HCMV gM Amino Acid Sequence

MAPSHVDKVNTRTWSASIVFMVLTFVNVSVHLVLSNFPHLGYPCVYYHVVDFE RLNMSAYNVMHLHTPMLFLDSVQLVCYAVFMQLVFLAVTIYYLVCWIKISMRK DKGMSLNQSTRDISYMGDSLTAFLFILSMDTFQLFTLTMSFRLPSMIAFMAAVHF FCLTIFNVSMVTQYRSYKRSLFFFSRLHPKLKGTVQFRTLIVNLVEVALGFNTTVV AMALCYGFGNNFFVRTGHMVLAVFVVYAIISIIYFLLIEAVFFQYVKVQFGYHLG AFFGLCGLIYPIVQYDTFLSNEYRTGISWSFGMLFFIWAMFTTCRAVRYFRGRGS GSVKYQALATASGEEVAVLSHHDSLESRRLREEEDDDDDDEDFEDA* (SEQ ID NO:4)

Matter No.: 07917-190001 Page 13 of 20 Applicant(s): Lu et al. COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

HCMV gN cDNA Sequence

ATGGAGTGGAACACTAGTATTAGGTCTTTTAGTTTTATCGGTAGTGGCAGA GAGTTCTGGTAACAACTACAACGTCAACCTCTGCAACTACAACAACTACAAAGTCTT CTGCTAGCGTATCAACTACCAAACTAACAACAACTGCAACAACTACCAACAACTACGACGACTACGACCTAATCGACAACTAGCACTAAACTCAGTTCTACC ACCCACGATCCTAATGTGATGAGACGACATGCGAACGATGATTTTTACAAGG CGCATTGCACATCGCATATGTATGAGGCTCTCACTGTCCAGCTTTGCGGCCTGG TGGACTATGCTTAATGCTCTAATTCTCATGGGAGCTTTTTGTATTGTACTACG ACATTGCTGCTCCAGAACTTTACTGCAACCACCACCAAAGGCTATTGA (SEQ ID NO:5)

HCMV gN Amino Acid Sequence

MEWNTLVLGLLVLSVVAESSGNNSSTSTSATTSKSSASVSTTKLTTVATTSATTT TTTTLSTTSTKLSSTTHDPNVMRRHANDDFYKAHCTSHMYELSLSSFAAWWTM LNALILMGAFCIVLRHCCFQNFTATTTKGY* (SEQ ID NO:6)

Page 14 of 20

Matter No.: 07917-190001 Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

TREATMENT

HCMV gH cDNA Sequence

ATGCGGCCGGCCTCCCCCCTACCTCACTGTCTTCACCGTCTACCTCCTCAGTCACC TACCTTCGCAACGATATGGCGCGGACGCCGCATCCGAAGCGCTGGACCCTCACGCAT TTCACCTACTACTCAACACCTACGGGAGACCCATCCGCTTCCTGCGTGAAAACACCA CCCAGTGCACCTACAACAGCAGCCTCCGTAACAGCACGGTCGTCAGGGAAAACGCC ATCAGTTTCAACTTTTTCCAAAGCTATAATCAATACTATGTATTCCATATGCCTCGAT GTCTTTTTGCGGGTCCTCTGGCGGAGCAGTTTCTGAACCAGGTAGATCTGACCGAAA CCCTAGAAAGATACCAACAGAGACTTAACACCTACGCATTGGTATCCAAAGACCTGG CCAGCTACCGATCTTTTCGCAGCAGCTGAAGGCACAAGACAGCCTGGGTCAGCAGC CCACCACCGTGCCACCGCCCATTGATCTGTCAATACCTCACGTTTGGATGCCACCCCA AACCACTCCACACGACTGGAAGGGATCGCACACCACCTCGGGACTACATCGGCCAC ACTTTAACCAGACCTGTATCCTCTTTGATGGACACGATCTGCTTTTCAGCACCGTTAC GCCCTGTCTGCACCAGGGCTTTTACCTCATGGACGAACTACGTTACGTTAAAATCAC ACTGACCGAGGACTTCTTCGTAGTTACGGTATCTATAGACGACGACACCCATGCT GCTTATCTTCGGTCATCTTCCACGCGTACTCTTCAAAGCGCCCTATCAACGCGACAAC TTTATACTACGACAAACTGAAAAACACGAGCTCCTGGTACTAGTTAAGAAAGCTCAA CTAAACCGTCACTCCTATCTCAAAGACTCGGACTTCTCGACGCCGCACTCGACTTCA ACTACCTGGACCTCAGCGCACTGTTACGTAACAGCTTTCACCGTTACGCTGTAGACGT ACTCAAAAGCGGTCGATGTCAAATGTTGGACCGCCGCACGGTAGAAATGGCCTTCGC CTACGCATTAGCACTGTTCGCGGCAGCCCGACAAGAAGAGGCCGGCACCGAAATCTC CATCCCACGAGCCCTAGACCGCCAGGCCGCACTCTTACAAATACAAGAATTTATGAT CACCTGCCTCTCACAAACACCACCACGCACCACATTGCTGCTATATCCCACAGCCGT GGACCTGGCCAAACGAGCCCTCTGGACGCCGGACCAGATCACCGACATCACCAGCCT CGTACGCCTGGTCTACATACTTTCTAAACAGAATCAGCAACATCTCATTCCCCAGTGG GCACTACGACAGATCGCCGACTTTGCCCTACAATTACACAAAACGCACCTGGCCTCT TTTCTTTCAGCCTTCGCGCGCCAAGAACTCTACCTCATGGGCAGCCTCGTCCACTCCA TGTTGGTACATACGACGGAGAGACGCGAAATCTTCATCGTAGAAACGGGCCTCTGTT CATTGGCCGAGCTATCACACTTTACGCAGTTGCTAGCTCATCCGCACCACGAATACCT CCTCACGCGTCTCTCCCGATGCCACCGTTCCTGCTACCGTTCCCGCCGCCCTCTCC ATCCTATCTACCATGCAACCAAGCACGCTGGAAACCTTCCCCGACCTGTTTTGTCTGC CGCTCGGCGAATCCTTCTCCGCGCTAACCGTCTCCGAACACGTCAGTTATGTCGTAAC AAACCAGTACCTGATCAAAGGTATCTCCTACCCTGTCTCCACCACCGTCGTAGGCCA GAGCCTCATCACCCAAACGGACAGTCAAACTAAATGCGAACTAACGCGCAACA TGCACACCACACAGCATCACAGCGGCGCTCAACATTTCACTAGAAAACTGCGCCT TTTGCCAAAGCGCCTGCTAGAATACGACGACACGCAAGGCGTCATCAACATCATGT ACATGCACGACTCGGACGTCCTTTTCGCCCTGGATCCCTACAACGAAGTGGTGG TCTCATCTCCGCGAACTCACTACCTCATGCTTTTGAAAAACGGTACGGTCCTAGAAGT AACTGACGTCGTCGTGGACGCCACCGACAGTCGTCTCCTCATGATGTCCGTCTACGC GCTATCGGCCATCATCGGCATCTATCTGCTCTACCGCATGCTCAAGACATGCTGA (SEQ ID NO:7)

Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

Page 15 of 20

TREATMENT

HCMV gH Amino Acid Sequence

MRPGLPPYLTVFTVYLLSHLPSQRYGADAASEALDPHAFHLLLNTYGRPIRFLRE NTTQCTYNSSLRNSTVVRENAISFNFFQSYNQYYVFHMPRCLFAGPLAEQFLNQV DLTETLERYQQRLNTYALVSKDLASYRSFSQQLKAQDSLGQQPTTVPPPIDLSIPH VWMPPQTTPHDWKGSHTTSGLHRPHFNQTCILFDGHDLLFSTVTPCLHQGFYLM DELRYVKITLTEDFFVVTVSIDDDTPMLLIFGHLPRVLFKAPYQRDNFILRQTEKH ELLVLVKKAQLNRHSYLKDSDFLDAALDFNYLDLSALLRNSFHRYAVDVLKSGR CQMLDRRTVEMAFAYALALFAAARQEEAGTEISIPRALDRQAALLQIQEFMITCL SQTPPRTTLLLYPTAVDLAKRALWTPDQITDITSLVRLVYILSKQNQQHLIPQWAL RQIADFALQLHKTHLASFLSAFARQELYLMGSLVHSMLVHTTERREIFIVETGLCS LAELSHFTQLLAHPHHEYLSDLYTPCSSSGRRDHSLERLTRLFPDATVPATVPAA LSILSTMQPSTLETFPDLFCLPLGESFSALTVSEHVSYVVTNQYLIKGISYPVSTTV VGQSLIITQTDSQTKCELTRNMHTTHSITAALNISLENCAFCQSALLEYDDTQGVI NIMYMHDSDDVLFALDPYNEVVVSSPRTHYLMLKKNGTVLEVTDVVVDATDSR LLMMSVYALSAIIGIYLLYRMLKTC* (SEQ ID NO:8)

Matter No.: 07917-190001 Page 16 of 20

Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

TREATMENT

HCMV gL cDNA Sequence

CGTTTTAGGGATCGAAGACCTGAGCGCCAACTTTCGGCGCCAACTGGCTCCTT ACCGTCACACTCTCATCGTGCCGCAGACTTGATGTGCCGCCGCCCGGATTGCG GCTTCTCTTCTCACCTGGACCGGTGGTACTGCTGTGGTGTTTCTCTGCTGC CCATTGTTTCCTCAGTCGCCGTCAGCGTCGCTCCTACCGCCGCCGAGAAAGTC CCCGCGGAGTGCCCCGAACTAACGCGTCGATGCCTGTTGGGTGAGGTGTTTC AGGGTGACAAGTATGAAAGTTGGCTGCGCCCGTTGGTGAATGTTACCAGACG CGATGGCCCGCTATCGCAACTTATTCGTTACCGTCCCGTTACGCCGGAGGCCG CCAACTCCGTGCTGTTGGACGATGCTTTCCTGGACACTCTGGCCCTGCTGTAC AACAATCCGGATCAATTGCGGGCCCTGCTGACGCTGTTGAGCTCGGACACAG CGCCGCGCTGGATGACGGTGATGCGCGGCTACAGCGAGTGCGGCGATGGCTC GCCGCCGTGTACACGTGCGTGGACGACCTGTGCCGCGGCTACGACCTCACG CGACTGTCATACGGCGCAGCATCTTCACGGAACACGTGTTAGGCTTCGAGC TGGTGCCACCGTCTCTCTTTAACGTGGTGGTGGCCATACGCAACGAAGCCAC GCGTACCAACCGCGCGTGCGTCTGCCCGTGAGCACCGCTGCCGCGCCCGAG TCACCAGCTGGACCCGCCGCTACTACGCCACCTAGATAAATACTACGCCGGA CTGCCGCCCGAGCTGAAGCAGACGCGCGTCAACCTGCCGGCTCACTCGCGCT ATGGCCCTCAAGCAGTGGATGCTCGCTAA (SEQ ID NO:9)

HCMV gL Amino Acid Sequence

RFRDRRPERQLSAPTGSLPSHSHRAADLMCRRPDCGFSFSPGPVVLLWCCLLLPI VSSVAVSVAPTAAEKVPAECPELTRRCLLGEVFQGDKYESWLRPLVNVTRRDGP LSQLIRYRPVTPEAANSVLLDDAFLDTLALLYNNPDQLRALLTLLSSDTAPRWMT VMRGYSECGDGSPAVYTCVDDLCRGYDLTRLSYGRSIFTEHVLGFELVPPSLFNV VVAIRNEATRTNRAVRLPVSTAAAPEGITLFYGLYNAVKEFCLRHQLDPPLLRHL DKYYAGLPPELKQTRVNLPAHSRYGPQAVDAR* (SEQ ID NO:10)

Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

Page 17 of 20

TREATMENT

HCMV gO cDNA Sequence

ATGGGGAGAAAAGAGATGATGGTGAGAGACGTCCCTAAGATGGTGTTTCTAA TATCTATATCTTCTTGCTTGTTTCTTCATAAACTGTAAAGTTATGTCAAAAG CGCTTTATAATCGTCCTTGGAGGGGCTTGGTACTGTCTAAGATAGGCAAATAT AAATTAGATCAGCTTAAGTTAGAAATTTTGAGACAACTAGAAACGACTATTT CTACAAAATACAATGTAAGTAAACAACCGGTTAAAAAATCTCACTATGAACAT GACAGAGTTTCCACAATACTACATTTTAGCGGGCCCCATTCAGAATTATAGTA TAACCTATCTGTGGTTTGATTTTTATAGTACCCAGCTTAGAAAACCCGCAAAA TACGTTTACTCACAGTACAATCATACGGCTAAAACGATAACATTCAGACCCC CACCTTGTGGTACTGTCCTTCCATGACTTGTCTTTCCGAAATGCTAAACGTTT CCAAACGTAATGATACTGGCGAACAAGGTTGCGGTAATTTCACCACGTTCAA CCCCATGTTTTCAATGTACCGCGTTGGAACACCAAATTGTACGTGGGTCCGA CTAAGGTTAACGTAGATAGTCAAACGATTTATTTTCTAGGTTTAACCGCCCTG CTTTTACGTTACGCACAACGCAACTGTACACACAGTTTCTACCTGGTTAACGC CATGAGCCGGAATCTATTTCGCGTCCCCAAGTATATTAACGGCACCAAGTTA AAAAACACTATGCGAAAACTAAAACGTAAACAAGCGCCCGTTAAGGAACAA TTCGAAAAAAAGCTAAGAAAACTCAGAGTACTACTACGCCATACTTTTCCT ATACAACGTCTGCCGCTCTCAACGTCACTACTAACGTGACTTATAGTATTACT ACCGCCGCAAGGCGGGTTTCCACGTCTACAATTGCTTATCGTCCTGATAGCAG CTTTATGAAGTCCATTATGGCCACACAGTTAAGGGACCTAGCAACGTGGGTG TATACCACTCTACGTTACCGGCAAAATCCTTTTTGTGAACCAAGCCGCAACCG AACCGCCGTGTCAGAATTTATGAAAAACACGCACGTACTAATCCGTAACGAA ACGCCGTACACTATTTACGGTACTCTCGACATGAGCTCCTTATATTACAACGA AACCATGTTCGTGGAAAACAAAACAGCTTCCGATAGTAACAAAACTACACCT ACGTCACCATCAATGGGGTTTCAGAGAACATTTATAGATCCCCTGTGGGACT ATCTAGACTCGCTGCTTTTCTAGATGAGATTCGTAACTTTAGCCTCCGGTCA CCCACGTATGTAAACCTTACCCCGCCGGAACACCCGCCGGGCTGTAAATCTGT CCACCCTCAATAGCCTTTGGTGGTGGTTGCAGTAA (SEO ID NO:11)

HCMV gO Amino Acid Sequence

MGRKEMMVRDVPKMVFLISISFLLVSFINCKVMSKALYNRPWRGLVLSKIGKYK LDQLKLEILRQLETTISTKYNVSKQPVKNLTMNMTEFPQYYILAGPIQNYSITYLW FDFYSTQLRKPAKYVYSQYNHTAKTITFRPPPCGTVPSMTCLSEMLNVSKRNDT GEQGCGNFTTFNPMFFNVPRWNTKLYVGPTKVNVDSQTIYFLGLTALLLRYAQR NCTHSFYLVNAMSRNLFRVPKYINGTKLKNTMRKLKRKQAPVKEQFEKKAKKT QSTTTPYFSYTTSAALNVTTNVTYSITTAARRVSTSTIAYRPDSSFMKSIMATQLR DLATWVYTTLRYRQNPFCEPSRNRTAVSEFMKNTHVLIRNETPYTIYGTLDMSSL YYNETMFVENKTASDSNKTTPTSPSMGFQRTFIDPLWDYLDSLLFLDEIRNFSLRS PTYVNLTPPEHRRAVNLSTLNSLWWWLQ* (SEQ ID NO:12)

Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

Page 18 of 20

TREATMENT

HCMV pp65 cDNA Sequence

ATGGAGTCGCGGGTCGCCGTTGTCCCGAAATGATATCCGTACTGGGTCCCATTTCG GGGCACGTGCTGAAAGCCGTGTTTAGTCGCGGCGATACGCCGGTGCTGCCGCACGAG GTATCGCAGTACACGCCCGACTCGACGCCATGCCACCGCGGCGACAATCAGCTGCAG GTGCAGCACGTACTTTACGGGCAGCGAGGTGGAGAACGTGTCGGTCAACGTGCA CAACCCCACGGCCGAAGCATCTGCCCCAGCCAGGAGCCCATGTCGATCTATGTGTA CGCGCTGCCGCTCAAGATGCTGAACATCCCCAGCATCAACGTGCACCACTACCCGTC GGCGGCCGAGCGCAAACACCGACACCTGCCCGTAGCTGACGCTGTGATTCACGCGTC GGGCAAGCAGATGTGGCAGGCGCGTCTCACGGTCTCGGGACTGGCCTGGACGCGTC AGCAGAACCAGTGGAAAGAGCCCGACGTCTACTACACGTCAGCGTTCGTGTTTCCCA CCAAGGACGTGCACTGCGCACGTGGTGTGCGCGCACGAGCTGGTTTGCTCCATGG AGAACACGCGCGCAACCAAGATGCAGGTGATAGGTGACCAGTACGTCAAGGTGTAC CTGGAGTCCTTCTGCGAGGACGTGCCCTCCGGCAAGCTCTTTATGCACGTCACGCTG GGCTCTGACGTGGAAGAGGACCTGACGATGACCCGCAACCCGCAACCCTTCATGCGC CCCCACGAGCGCAACGGCTTTACGGTGTTGTGTCCCAAAAATATGATAATCAAACCG GGCAAGATCTCGCACATCATGCTGGATGTGGCTTTTACCTCACACGAGCATTTTGGG CTGCTGTGTCCCAAGAGCATCCCGGGCCTGAGCATCTCAGGTAACCTGTTGATGAAC GGGCAGCAGATCTTCCTGGAGGTACAAGCCATACGCGAGACCGTGGAACTGCGTCA GTACGATCCCGTGGCTGCGCTCTTCTTTTTCGATATCGACTTGCTGCTGCAGCGCGGG CCTCAGTACAGCGAGCACCCCACCTTCACCAGCCAGTATCGCATCCAGGGCAAGCTT GAGTACCGACACCTGGGACCGCACGACGAGGGTGCCGCCCAGGGCGACGACGA CGTCTGGACCAGCGGATCGGACTCCGACGAGAACTCGTAACCACCGAGCGCAAGA CGCCCGCGTCACCGGCGCGCGCCCATGGCGGGCGCCTCCACTTCCGCGGGCCGCA AACGCAAATCAGCATCCTCGGCGACGGCGTGCACGTCGGGCGTTATGACACGCGGCC GCCTTAAGGCCGAGTCCACCGTCGCGCCCGAAGAGGACACCGACGAGGATTCCGAC AACGAAATCCACAATCCGGCCGTGTTCACCTGGCCGCCCTGGCAGGCCGGCATCCTG GCCCGCAACCTGGTGCCCATGGTGGCTACGGTTCAGGGTCAGAATCTGAAGTACCAG GAATTCTTCTGGGACGCCAACGACATCTACCGCATCTTCGCCGAATTGGAAGGCGTA TGGCAGCCCGCTGCGCAACCCAAACGTCGCCGCCACCGGCAAGACGCCTTGCCCGGG CCATGCATCGCCTCGACGCCCAAAAAGCACCGAGGTTGA (SEO ID NO:13)

HCMV pp65 Amino Acid Sequence

MESRGRRCPEMISVLGPISGHVLKAVFSRGDTPVLPHETRLLQTGIHVRVSQPSLILVSQY TPDSTPCHRGDNQLQVQHTYFTGSEVENVSVNVHNPTGRSICPSQEPMSIYVYALPLKML NIPSINVHHYPSAAERKHRHLPVADAVIHASGKQMWQARLTVSGLAWTRQQNQWKEPD VYYTSAFVFPTKDVALRHVVCAHELVCSMENTRATKMQVIGDQYVKVYLESFCEDVPS GKLFMHVTLGSDVEEDLTMTRNPQPFMRPHERNGFTVLCPKNMIIKPGKISHIMLDVAFT SHEHFGLLCPKSIPGLSISGNLLMNGQQIFLEVQAIRETVELRQYDPVAALFFFDIDLLLQR GPQYSEHPTFTSQYRIQGKLEYRHTWDRHDEGAAQGDDDVWTSGSDSDEELVTTERKT PRVTGGGAMAGASTSAGRKRKSASSATACTSGVMTRGRLKAESTVAPEEDTDEDSDNEI HNPAVFTWPPWQAGILARNLVPMVATVQGQNLKYQEFFWDANDIYRIFAELEGVWQPA AQPKRRHRQDALPGPCIASTPKKHRG (SEQ ID NO:14)

Page 19 of 20

Matter No.: 07917-190001 Applicant(s): Lu et al.

COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS

TREATMENT

HCMV pp150 cDNA Sequence

ATGAGTTTGCAGTTTATCGGTCTACAGCGGCGCGATGTGGTAGCCCTGGTCAACTTTCTGCGC CATCTCACGCAAAAGCCCGACGTGGATCTCGAGGCACACCCCAAGATCCTGAAAAAATGTGG CCGCGAGCTGCGTTTTCACAACCCCGACCTCTCCTCAGTGCTCGAGGAGTTCGAGGTGCGTTG CGTGGCCGTGGCGCGCGCGCTACACTTACCCGTTCGGTGATCGTGGTAAGGCGCGTGACCA CCTGGCTGTGCTAGACCGTACCGAATTCGATACGGACGTGCGCCACGATGCCGAGATCGTGGA ACGCGCGCTCGTAAGCGCGGTCATTCTGGCCAAGATGTCGGTGCGCGAGACGCTGGTCACAG CCATCGGCCAGACGGAACCCATCGCCTTTGTGCACCTCAAGGATACGGAGGTGCAGCGCATTG AAGAAAACCTGGAGGTGTGCGCCGTAACATGTTCTGCGTGAAACCGCTCGACCTTAACCTGG ACCGGCACGCCAACACGGCGCTGGTCAACGCCGTCAACAAGCTCGTGTACACGGGCCGTCTC ATCATGAACGTGCGCAGGTCTTGGGAGGAGCTGGAGCGCAAATGTCTGGCGCGCATTCAGGA GCGCTGCAAGCTGCTCAAGGAGCTGCGCATGTGCCTTTCCTTTGATTCCAACTACTGTCGC AATATCCTCAAGCACGCCGTGGAAAACGGCGACTCGGCCGACACGCTGTTGGAGCTGCTCATC CGTCGGTACCCAAGAAACATGTCCCCACGCAGCCGCTGGACGGCTGGAGCTGGATCGCCAGT CCCTGGAAGGGACACAAACCGTTCCGCTTCGAGGCCCATGGTTCTCTGGCACCGGCCGCGAA GCCCACGCTGCCCGTTCGGCGCCGTCGGCTATTACGACGAAGAGGAAAAGCGTCGCGAGCG GCAGAAACGGGTGGACGACGAGGTGGTGCAGCGTGAGAACAGCAGCTGAAGGCTTGGGAG GAGAGGCAGCAGCACCTGCAGCAACGTCAGCAGCACCCCCCGGCACGTAAACCGAG CGCCTCCGGAGGCTCTTTGGCTCCAGTGCCGATGAGGACGACGACGATGATGATGACGAGA AAAACATCTTTACGCCCATCAAGAAACCGGGAACTAGCGGCAAGGGCGCCGCTAGTGGTGGC GGTGTTTCCAGCATTTTCAGCGGCCTGTTATCCTCGGGCAGTCAGAAACCGACCAGCGGTCCC TTGAACATCCCGCAACAACAACAGCGTCACGCGGCTTTCAGTCTCGTCTCCCCGCAGGTGACC AAGGCCAGCCGGGAAGGGTCCGTCGGGACAGCGCGTGGGACGTGAGGCCGCTCACGGAGA CCAGAGGGGATCTTTTCTCGGGCGACGAGGATTCCGACAGCTCGGATGGCTATCCCCCAACC GTCAAGATCCGCGTTTCACCGACACGCTGGTGGACATCACGGATACCGAGACGAGCGCCAAA CCGCCCGTCACCACCGCGTACAAGTTCGAGCAACCGACGTTGACGTTCGGCGCCGGAGTTAAC GTTCCTGCTGGCGCCGGCGCTGCCATCCTCACGCCGACGCCTGTCAATCCTTCCACGGCCCCC GCTCCGGCCCGACACCTACCTTCGCGGGTACCCAAACCCCGGTCAACGGTAACTCGCCCTGG GCTCCGACGCGCCGTTGCCCGGGGATATGAACCCCGCCAACTGGCCGCGCAACGCGCGTG GGCCCTCAGGATCCTCACCTGGCTTACAATCCCTTCAGGATGCCTACGACTTCCACGGCTTCT AGCGTCTCGGGACGCCGCTGATGAGGTTTGGGCTTTAAGGGACCAAACTGCAGAGTCACCGG TCGAAGACAGCGAGGAGGACGACTCCTCGGACACCGGCTCCGTCAGCCTGGGA CACACACACCGTCGTCCGATTACAACAACGACGTCATTTCGCCTCCCAGTCAGACGCCCGAG CAGTCGACGCCGTCCAGAATACGTAAAGCTAAGTTATCGTCTCCAATGACGACGACATCCACG AGCCAGAAACCGGTGCTGGGCAAGCGAGTCGCGACGCGCACGCGTCCGCCCGAGCGCAGAC GGTGACGTCGACGCCGGTTCAGGGAAGGCTAGAGAAACAGGTGTCGGGCACGCCGTCGACGG TACCCGCCACGCTGTTGCAACCTCAACCGGCTTCGTCTAAAACGACGTCATCAAGGAACGTGA CTTCTGGCGCGGGAACCTCTTCCGCTTCTTCGGCTCGACAGCCGTCAGCCTCGGCGTCCGTTTT GTCGCCCACGGAGGATGATGTCGTGTCCCCCGCCACATCGCCGCTGTCCATGCTTTCGTCAGC CTCTCCGTCCCGGCCAAGAGTGCCCCCCGTCTCCGGTGAAAGGCCGGGGCAGCCGCGTCGG TGTTCCTTCAAACCTACTTTGGGCGGCAAGGCGGTGGTAGGTCGACCGCCCTCGGTCCC CGTGAGCGGTAGCGCCGGGTCGCCTGTCCGGCAGCAGCCGGGCCGCCTCGACCACGCCGA CGTATCCCGCGGTAACCACCGTTTACCCACCGTCGTCTACGGCCAAAAGCAGCGTATCGAATG GCTCGACGGGGACCGCCGCGTAGGTTCCCCCGTCAAGAGCACGACGGGCATGAAAACGGTG GCTTTCGACCTATCGTCGCCCCAGAAGAGCGGTACGGGGCCGCAACCGGGTTCTGCCGGCATG GGGGCCCAAAACGCCGTCGGACGCCGTGCAGAACATCCTCCAAAAGATCGAGAAGATTAA GAACACGGAGGAATAG (SEQ ID NO:15)

Matter No.: 07917-190001 Page 20 of 20 Applicant(s): Lu et al. COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS TREATMENT

HCMV pp150 Amino Acid Sequence

MSLQFIGLQRRDVVALVNFLRHLTQKPDVDLEAHPKILKKCGEKRLHRRTVLFN ELMLWLGYYRELRFHNPDLSSVLEEFEVRCVAVARRGYTYPFGDRGKARDHLA VLDRTEFDTDVRHDAEIVERALVSAVILAKMSVRETLVTAIGQTEPIAFVHLKDT EVORIEENLEGVRRNMFCVKPLDLNLDRHANTALVNAVNKLVYTGRLIMNVRR SWEELERKCLARIQERCKLLVKELRMCLSFDSNYCRNILKHAVENGDSADTLLEL LIEDFDIYVDSFPQSAHTFLGARSPSLEFDDDANLLSLGGGSAFSSVPKKHVPTQP LDGWSWIASPWKGHKPFRFEAHGSLAPAAEAHAARSAAVGYYDEEEKRREROK RVDDEVVQREKQQLKAWEERQQNLQQRQQQPPPPARKPSASRRLFGSSADEDD DDDDDEKNIFTPIKKPGTSGKGAASGGGVSSIFSGLLSSGSQKPTSGPLNIPQQQQ RHAAFSLVSPOVTKASPGRVRRDSAWDVRPLTETRGDLFSGDEDSDSSDGYPPN RQDPRFTDTLVDITDTETSAKPPVTTAYKFEQPTLTFGAGVNVPAGAGAAILTPTP VNPSTAPAPAPTPTFAGTQTPVNGNSPWAPTAPLPGDMNPANWPRERAWALKN PHLAYNPFRMPTTSTASQNTVSTTPRRPSTPRAAVTQTASRDAADEVWALRDQT AESPVEDSEEEDDDSSDTGSVVSLGHTTPSSDYNNDVISPPSQTPEQSTPSRIRKAK LSSPMTTTSTSQKPVLGKRVATPHASARAQTVTSTPVQGRLEKQVSGTPSTVPAT LLOPOPASSKTTSSRNVTSGAGTSSASSAROPSASASVLSPTEDDVVSPATSPLSM LSSASPSPAKSAPPSPVKGRGSRVGVPSLKPTLGGKAVVGRPPSVPVSGSAPGRLS GSSRAASTTPTYPAVTTVYPPSSTAKSSVSNAPPVASPSILKPGASAALQSRRSTG TAAVGSPVKSTTGMKTVAFDLSSPQKSGTGPQPGSAGMGGAKTPSDAVQNILQK **IEKIKNTEE (SEQ ID NO:16)**